

LANL Oralgen Questionnaire

===== **Part I: Oral Microbiome Sequencing Support:**

Please rank your priorities based on following list for your metagenome study? In order better for us to help, you are encouraged to tell us more about your microbiome project (of course all information received will be kept strictly confidential).

1) Metagenome Community Composition Profiling

e.g. determining whether there is a core metagenome of the human dental plaque from 16s rRNA surveys

2) Metagenome Sequence Processing

e.g. metagenome binning and assembly

3) Metadata Annotation

e.g. automatic annotation on GO terms, COGs, pathway assignments

4) Gene-centric Analyses

e.g. determining common/shared domains/genes/species/pathways between samples

Part II: Oralgen Web Site (<http://oralgen.lanl.gov/>):

Suggestion of improvements to be made

If you have any other comments or suggestions, please let us know.

1) Is the website interface intuitive for users? If not, what can we do to improve the user interface? What quicklinks or features could be added to enhance ease of use?

2) Are there any new tools that you would like to see implemented by Oralgen, in addition to the tools currently available?

3) We periodically include genome-specific analyses for some of the genomes on the Oralgen website. If you would like to see specialized analyses for any particular Oralgen genome, could you please describe?